

Db	3075	GGTTTGAATAATTGCGAGGCCGCGCATGATCTTCTCACTTACTCTCTCCCAATTC	3334
Qy	1448	AGCAATCAACCCAGACTAAGAGGATTTTGTATCCCTAGTGATTACAGCCCTCAAGAAATCAA	1507
Db	3135	AGCATTCACACCAAGACTAAGAGGATTTTGTATCCCTAGTGATTACAGCCCTGAGGAAATCAA	3194
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Qy	1748	CCCTCAACCGTTGTGTACTGACCATTTTCCACTACATGCCCTTCCCTACCTTCCCTTCACAA	1807
Db	3435	CCCTCAACCGTTGTGTACTGACCATTTTCCACTACATGCCCTTCCCTACCTTCCCTTCACAA	3494
Qy	1808	CCCAATCAAGTGAATACCTTGATTAATATCTCTCTTACTGTGCTTATCTTTTCTTTGTTG	1867
Db	3495	CCCAATCAAGTGAATACCTTGATTAATATCTCTCTTACTGTGCTTATCTTTTCTTTGTTG	3554
Qy	1868	GATTGGTTCTAAATTAAATGAATAAAGATTTCTAAATTTACATTTTATAGGGTATTGTA	1927
Db	3555	GATTGGTTCTAAATTAAATGAATAAAGATTTCTAAATTTACATTTTATAGGGTATTGTA	3614
Qy	1928	AATAAAACAAATGTATCTT	1949
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[illegible]

30-MAY-1997 US 60/048357, 30-MAY-1997 US 60/048189 PR
06-JUN-1997 US 60/048970, 05-SEP-1997 US 60/057765 PR
19-DEC-1997 US 69/068368
PI STEVEN M RUBEN, CRAIG A ROSEN, YI LI, ZHIZHEN ZENG, HLA KYAW, PI
CARRIE L FISCHER,
PI HAODONG LI, DANIEL R SOPPET, REINER L GENTZ, YING FEI WEI, PAUL A
PI MOORE,
PI PAUL E YOUNG, JOHN M GREENE, ANN M FERRIE
PC C12N15/12, C07K14/47, C12Q1/68, C12N1/21, C12N15/62, C07K19/40, PC
C12N5/10, C07K16/18, G01N33/60, G01N33/68, C12Q1/48, G01N33/53, A61K38/17, PC
A61K48/00, C12N15/11, C12N1/21, C12R1/19)
PC A61K31/70, C12N15/11, C12N1/21, C12N15/62, C07K19/40, PC
CC Strandedness: Double;
CC Topology: Linear;
CC 28 human secreted proteins
FH Key Location/Qualifiers
FT source i. .2350 /organism="Unidentified"
FT Location/Qualifiers
1. .2350
/organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
BASE COUNT 612 575 C 419 G 737 T 7 others
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Query Match 97.0%; Score 1914.2; DB 6; Length 2350;
Best Local Similarity 99.5%; Pred. No. 0;
Matches 1943; Conservative 7; Mismatches 0; Indels 3; Gaps 3;
QY 9 CAGAACAGTTGGCTGTCACCTGCTCTCATCTCTGAGCCGCTGCTGAGACCTACAG 68
Db 398 CAGAACAGTTGGCTGTCACCTGCTCTCATCTCTGAGCCGCTGCTGAGACCTACAG 457
QY 69 AGTTCAGACCTGCTGAGAGTCAACACAAATTAATCAGTGGAGCGAGGTGATGTTGATG 128
Db 458 AGTTCAGACCTGCTGAGAGTCAACACAAATTAATCAGTGGAGCGAGGTGATGTTGATG 517
QY 129 GCAACACGAGAACTTCTGCGAATCCCAAGCAGCCCTGCGCAAGCCCAATCTCTATACCT 188
Db 518 GCAACACGAGAACTTCTGCGAATCCCAAGCAGCCCTGCGCAAGCCCAATCTCTATACCT 577
QY 189 ACAACCTAGTGTCTCTCTTTTCTGCTATGTTGACCGGACAGAACTGAACTGCTGGAG 248
Db 578 ACAACCTAGTGTCTCTCTTTTCTGCTATGTTGACCGGACAGAACTGAACTGCTGGAG 637
QY 249 ACAACCTAGTGTCTCTCTTTTCTGCTATGTTGACCGGACAGAACTGAACTGCTGGAG 308
Db 638 ASAAACCTGACGCTAGTTCAGGGAGTACATAAAGATCTCCAAATGTGCTGCTACCTC 697
QY 309 CTTTGTGCTGCACTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 368
Db 698 CTTTGTGCTGCACTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 757
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Db 758 AGCTAGCTTTTCCCAAGCAAGTCTTGTGGGATCGAATTCCTGTTTACAGGGGTTGTC 817
QY 429 TCTCTAAATGTCAGATATTTCCCAAGCAAGTCTTGTGGGATCGAATTCCTGTTTACAGGGGTTGTC 488
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QY

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1268 ATGTAACCAATGAAGCTAAGTTGGCTGCTGCTTTTGTAGTCTCCACACCTAGGGCAG 1327
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2318 TAAATAAAAACAATTTGATATCTTAAATAAAAA 2350

PAT 27-AUG-2002

BD084164 2350 bp DNA linear
NITION 28 human secreted proteins.
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ION BD084164.1 GI:22629774
ORDS JP 2001524814-A/12.
CE unclassified
GANISM unclassified.
RENCE 1 (bases 1 to 2350)
THORS Ruben, S.M., Rosen, C.A., Li, Y., Zeng, Z., Kyaw, H., Fisch, C.L., Li, H.,
Soppet, D.R., Gentz, R.L., Wei, Y., Moore, P.A., Young, P.E.,
Greene, J.M. and Ferrie, A.M.
28 human secreted proteins
HUMAN GENOME SCIENCES INC
OS Unidentified
PD JP 2001524814-A/12
PF 04-DEC-2001
PR 12-MAR-1998 JP 1998539800
PR 14-MAR-1997 US 60/040762, 14-MAR-1997 US 60/048210 PR
30-MAY-1997 US 60/050934, 30-MAY-1997 US 60/048100 PR